

FIG. 1

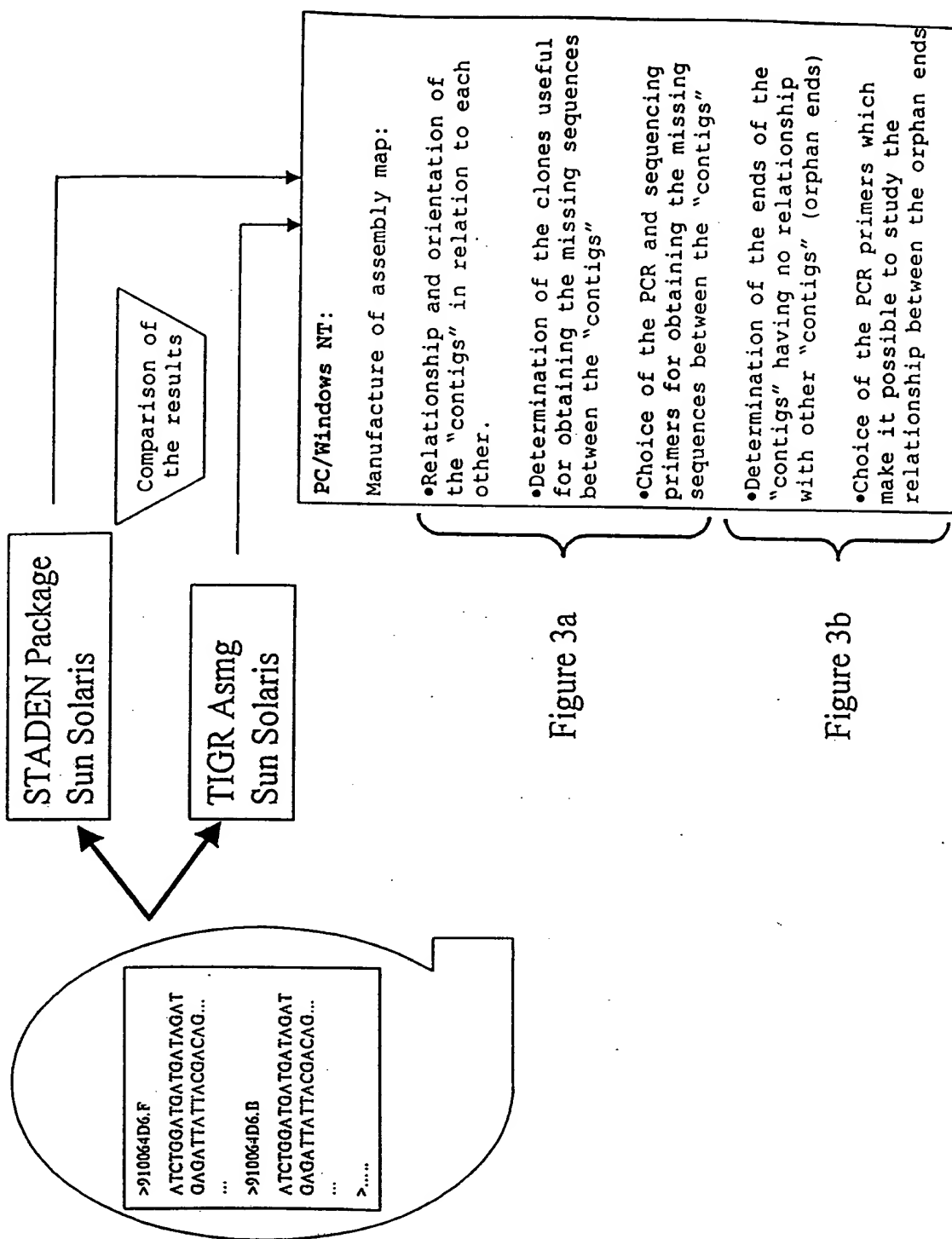


Figure 3a

Figure 3b

FIG. 2

a.

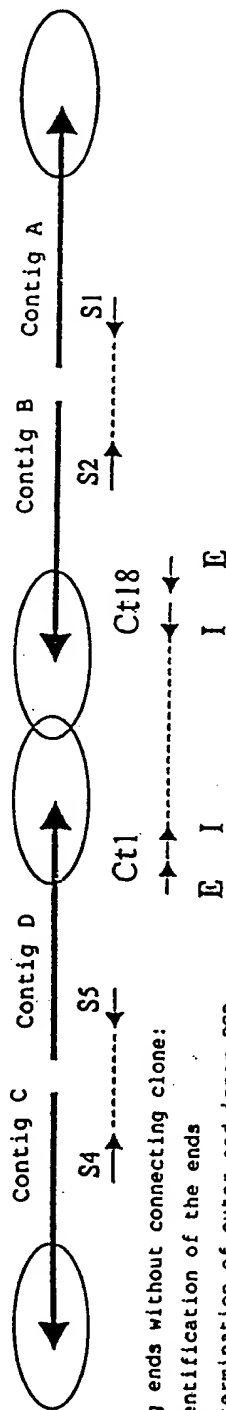


Distribution and position of the sequences on the contigs:
1- Define the arrangement of the contigs in relation to each other
2- Define the PCR primers which make it possible to fill the sequence

Statistical determination of the sequences:

- 1- Belonging to the same clone
- 2- Situated on two different contigs

b.



Contig ends without connecting clone:

1- Identification of the ends

2- Determination of outer and inner PCR

primers for studying the relationships

between the contigs

E: outer primers

I: inner primers

FIG. 3